

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Application of: Kessler, et al.



JUN 27 2002

Serial No.: 09/530,746

Group Art Unit: 1654

Filed: November 16, 2000

Examiner: Strzelecka, T.

For: SPECIFIC AND SENSITIVE METHOD
FOR DETECTING NUCLEIC ACIDS

Attorney Docket No.: 1803-277-999

AMENDMENT UNDER 37 C.F.R. § 1.111

Commissioner for Patents
Washington, D.C. 20231

Sir:

In response to the Notice to Comply with the Sequence Rules mailed September 7, 2001, please amend the above-identified application. This amendment is accompanied by (1) Appendix A showing a marked-up version of the amended claim; (2) Appendix B showing a clean copy of the claims as pending after entry of this amendment; (3) a Petition Pursuant to 37 C.F.R. § 1.137(b) to revive an unintentionally abandoned patent application; and (4) a Sequence Listing in compliance with 37 C.F.R. §§ 1.821-1.825 in paper and computer readable format.

AMENDMENT

IN THE CLAIMS:

Please amend the claim to read as follows:

22. (Twice Amended) A method for the detection of HCV, wherein two primers and one probe are used whose sequences are derived from sequences of consecutive bases the HGBV sequence shown below, complementary sequences thereto or sequences that are more than 80% identical to these sequences:

5'-GTACTGCCTGATAGGGCCTTGCAGGGGATCTGGAGTC
TCGTAGACCG TAGCACATG-3' (SEQ ID NO: 95).

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